

Exhibit 1

Translation of BM460207, 5'3' Frame 1

RTRGStopVSKHRWIPALRAWRLDGVSWRGKSGEAGTGIHRAICKGKStopMGStopRSYFGVS
 RQAPVSSStopSFRSVRLKNDPVLNLQKFSYTSSEDAWKTYLENPLTAATKAMMRVNGDDDSVA
 ALSFLYDYDMGPKERILSSSTGGRNDQGKRYHGMETDPLTSPHLMKFLTENVSGTPEY
 PDLLKKNLMSFGGLAHPWQSSPPCKPQQAGRLCGQVPVTHHStopVYVStopRPPSP
 PLFAKHFMGLPFNPAAGQPEPPLLSStopKNPTGDPSSLPPQYPCKTPPVNPLCSTKELHPTPPTRY
 ILPLNSLRPPPSKGIPPLPGPTAStopTPSSGSLTSHTRPPFLSLVSPSPSRETPL

Sequence Alignment

SEQ ID NO: 8	MWMNSILPIFLFRSVRLKNDPVLNLQKFSYTSSEDAWKTYLENPLTAATKAMMRVNGDDD	60
BM460207	-----SFRSVRLKNDPVLNLQKFSYTSSEDAWKTYLENPLTAATKAMMRVNGDDD	50

SEQ ID NO: 8	SVAALSFLYDYDMGPKERILSSSTGGRNDQGKRYHGMETDPLTSPHLMKXAAAL	120
BM460207	SVAALSFLYDYDMGPKERILSSSTGGRNDQGKRYHGMETDPLTSPHLMK--FL	108
	*****	*
SEQ ID NO: 8	TENVSGTPEYPDLLKKNLMSLEGALPTPGKAAPLPAGPSKLEAGSVDSYLLPXAATDMY	180
BM460207	TENVSGTPEYPDLLKKNLMSFGGLAHP-----	137
	*****; *, *	
SEQ ID NO: 8	DNGSLNSLFESIHGVPPTQRWQPDSTFKDDPOESMLFPDILKTSPEPPCEDYPSLKSDF	240
BM460207	-----WQSS-----PPCKPQ-----	149
	** , * ;	
SEQ ID NO: 8	EYTLGSPKAIHIKSGESPMAYLNKGQFYFVTLRTPAGGKGLALSSNKVKSVMVVFNEK	300
BM460207	-----AGGR-----LCG-----Q	157
	***: * , :	
SEQ ID NO: 8	VPVEQLRFWKHHWSRQPTAKQRVIDVADCKENFNTVEHIEEAVYALS FVWNVNBEAKVF	360
BM460207	VPVTH-----	163
	*** ;	
SEQ ID NO: 8	IGVNCCLSTDFSSQKGVKGVPLNLQIDTYDCGLGTERLVHRAVCQIKIFCDKGAERKMRDD	420
BM460207	-----	
SEQ ID NO: 8	ERKQFRKVKPCPDSSNSGVKGCLSGFRGNETTYLRPETDLETPPVLFIPNVHFSSLQRS	480
BM460207	-----	
SEQ ID NO: 8	GGAAPSAGPSSSNRLPLKRTCSPTTEBFEPLSPKQAKEGDLQRVLLYVRRETEEVFDALM	540
BM460207	-----	
SEQ ID NO: 8	LKTPDLKGLRNAISEKYGPPEENIYKVYKKCKRGILVNMDNNIIQHYSNHVAFLLDMGEL	600
BM460207	-----	
SEQ ID NO: 8	DGKIQIILKEL	611
BM460207	-----	